

SEQUENCE LISTING



110> Northwest Biotherapeutics, Inc.

<120> METHODS FOR THE DIAGNOSIS AND TREATMENT OF METASTATIC PROSTATE TUMORS

<130> 8511-025-228

<140> PCT/US99/08079

<141> 1999-04-13

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<170> PatentIn Ver. 2.0

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Leu Trp Leu Cys Leu Gly Leu Leu Asp Gly Leu Val Ser Asp Tyr Ser
15 20 25

atg acc ccc ccg acc ttg aac atc acg gag gag tca cac gtc atc gac

Met Thr Pro Pro Thr Leu Asn Ile Thr Glu Glu Ser His Val Ile Asp

30 35 40

acc ggt gac agc ctg tcc atc tcc tgc agg gga cag cac ccc ctc gag 195
Thr Gly Asp Ser Leu Ser Ile Ser Cys Arg Gly Gln His Pro Leu Glu
45 50 55

tgg gct tgg cca gga gct cag gag gcg cca gcc acc gga gac aag gac 243
Trp Ala Trp Pro Gly Ala Gln Glu Ala Pro Ala Thr Gly Asp Lys Asp

60 65 70 agc gag gac acg ggg gtg gtg cga gac tgc gag ggc aca gac gcc agg 291

age gag gae acg ggg gtg gtg ega gae tge gag gge aca gae gee agg 291 Ser Glu Asp Thr Gly Val Val Arg Asp Cys Glu Gly Thr Asp Ala Arg 75 80 85 90

ccc tac tgc aag gtg ttg ctg ctg cac gag gta cat gcc aac gac aca

Pro Tyr Cys Lys Val Leu Leu Leu His Glu Val His Ala Asn Asp Thr

95

100

105

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					gac Asp											483
			_	_	gtg Val 160							_	_	_	-	531
_		_	_		ctg Leu											579
					ctc Leu											627
					acc Thr											675
					atc Ile											723
_			_	_	ctg Leu 240		_	_	_			_	_	_		771
					gct Ala											819
				aag	cag	qca	gag						CCC	nan	003	~ ~ ~
	-	FIO	270	Lys	Gln											867
_	tcc	caa	270 cag	acc	Gln cac His	Ala aca	Glu gaa	Arg 275 ctc	Gly tcc	Lys agc	Trp atc	Val ctg	Pro 280 acc	Glu atc	Arg cac	915
Arg	tcc Ser	caa Gln 285 agc	270 cag Gln	acc Thr	cac	Ala aca Thr	Glu gaa Glu 290 ggc	Arg 275 ctc Leu tcg	Gly tcc Ser	Lys agc Ser	Trp atc Ile tgc	val ctg Leu 295	Pro 280 acc Thr	Glu atc Ile aac	cac His	
Arg aac Asn	tcc Ser gtc Val 300	caa Gln 285 agc Ser	cag Gln cag Gln	acc Thr cac His	cac His	aca Thr ctg Leu 305	gaa Glu 290 ggc Gly	Arg 275 ctc Leu tcg Ser	tcc ser tat Tyr	agc Ser gtg Val	atc Ile tgc Cys 310	ctg Leu 295 aag Lys	Pro 280 acc Thr gcc Ala	atc Ile aac Asn	cac His aac Asn	915

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	ccc Pro															1155
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	acc															1251
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	gag Glu															1347
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	tgg Trp 460															1443
	cgg Arg															1491
	gtg Val		_	-	_	_						_				1539
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	cag Gln															1635
	gtg Val 540															1683
	gac Asp															1731
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						ctc Leu										1827
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						gac Asp										2019
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Gln Ala Asp Ala			cag cgc cac agc ctg Gln Arg His Ser Leu 1225	3699
	Tyr Asn Trp Val		ggg tgc ctg gcc aga Gly Cys Leu Ala Arg 1240	3747

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Ile Ser Cys Arg Gly Gln His Pro Leu Glu Trp Ala Trp Pro Gly Ala
50 55 60

Gln Glu Ala Pro Ala Thr Gly Asp Lys Asp Ser Glu Asp Thr Gly Val
65 70 75 80

Val Arg Asp Cys Glu Gly Thr Asp Ala Arg Pro Tyr Cys Lys Val Leu 85 90 95

Leu Leu His Glu Val His Ala Asn Asp Thr Gly Ser Tyr Val Cys Tyr
100 105 110

Tyr	Lys	Tyr 115	Ile	Lys	Ala	Arg	Ile 120	Glu	Gly	Thr	Thr	Ala 125	Ala	Ser	Ser
Tyr	Val 130	Phe	Val	Arg	Asp	Phe 135	Glu	Gln	Pro	Phe	Ile 140	Asn	Lys	Pro	Asp
Thr 145	Leu	Leu	Val	Asn	Arg 150	Lys	Asp	Ala	Met	Trp 155	Val	Pro	Cys	Leu	Val 160
Ser	Ile	Pro	GÌу	Leu 165	Asn	Val	Thr	Leu	Arg 170	Ser	Gln	Ser	Ser	Val 175	Leu
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Val	Ser	Thr 195	Pro	Leu	Leu	His	Asp 200	Ala	Leu	Tyr	Leu	Gln 205	Cys	Glu	Thr
Thr	Trp 210	Gly	Asp	Gln	Asp	Phe 215	Leu	Ser	Asn	Pro	Phe 220	Leu	Val	His	Ile
Thr 225	Gly	Asn	Glu	Leu	Tyr 230	Asp	Ile	Gln	Leu	Leu 235	Pro	Arg	Lys	Ser	Leu 240
Glu	Leu	Leu	Val	Gly 245	Glu	Lys	Leu	Val	Leu 250	Asn	Cys	Thr	Val	Trp 255	Ala
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- Val Val Asn Val Pro Pro Gln Ile His Glu Lys Glu Ala Ser Ser Pro 420 425 430
- Ser Ile Tyr Ser Arg His Ser Arg Gln Ala Leu Thr Cys Thr Ala Tyr
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- Gly Val Pro Leu Pro Leu Ser Ile Gln Trp His Trp Arg Pro Trp Thr 450 455 460
- Pro Cys Lys Met Phe Ala Gln Arg Ser Leu Arg Arg Arg Gln Gln 465 470 475 480
- Asp Leu Met Pro Gln Cys Arg Asp Trp Arg Ala Val Thr Thr Gln Asp 485 490 495
- Ala Val Asn Pro Ile Glu Ser Leu Asp Thr Trp Thr Glu Phe Val Glu 500 505 510
- Gly Lys Asn Lys Thr Val Ser Lys Leu Val Ile Gln Asn Ala Asn Val 515 520 525
- Ser Ala Met Tyr Lys Cys Val Val Ser Asn Lys Val Gly Gln Asp Glu 530 540
- Arg Leu Ile Tyr Phe Tyr Val Thr Thr Ile Pro Asp Gly Phe Thr Ile 545 550 555 560
- Glu Ser Lys Pro Ser Glu Glu Leu Leu Glu Gly Gln Pro Val Leu Leu 565 570 575
- Ser Cys Gln Ala Asp Ser Tyr Lys Tyr Glu His Leu Arg Trp Tyr Arg 580 585 590
- Leu Asn Leu Ser Thr Leu His Asp Ala His Gly Asn Pro Leu Leu Leu 595 600 605
- Asp Cys Lys Asn Val His Leu Phe Ala Thr Pro Leu Ala Ala Ser Leu 610 620
- Glu Glu Val Ala Pro Gly Ala Arg His Ala Thr Leu Ser Leu Ser Ile 625 630 635 640
- Pro Arg Val Ala Pro Glu His Glu Gly His Tyr Val Cys Glu Val Gln 645 650 655
- Asp Arg Arg Ser His Asp Lys His Cys His Lys Lys Tyr Leu Ser Val 660 665 670
- Gln Ala Leu Glu Ala Pro Arg Leu Thr Gln Asn Leu Thr Asp Leu Leu 675 680 685
- Val Asn Val Ser Asp Ser Leu Glu Met Gln Cys Leu Val Ala Gly Ala 690 695 700
- His Ala Pro Ser Ile Val Trp Tyr Lys Asp Glu Arg Leu Leu Glu Glu 705 710 715 720

- Lys Ser Gly Val Asp Leu Ala Asp Ser Asn Gln Lys Leu Ser Ile Gln 725 730 735
- Arg Val Arg Glu Glu Asp Ala Gly Pro Tyr Leu Cys Ser Val Cys Arg
 740 745 750
- Pro Lys Gly Cys Val Asn Ser Ser Ala Ser Val Ala Val Glu Gly Ser 755 760 765
- Glu Asp Lys Gly Ser Met Glu Ile Val Ile Leu Val Gly Thr Gly Val
 770 780
- Ile Ala Val Phe Phe Trp Val Leu Leu Leu Leu Ile Phe Cys Asn Met 785 790 795 800
- Arg Arg Pro Ala His Ala Asp Ile Lys Thr Gly Tyr Leu Ser Ile Ile 805 810 815 `
- Met Asp Pro Gly Glu Val Pro Leu Glu Glu Gln Cys Glu Tyr Leu Ser 820 825 830
- Tyr Asp Ala Ser Gln Trp Glu Phe Pro Arg Glu Arg Leu His Leu Gly 835 840 845
- Arg Val Leu Gly Tyr Gly Ala Phe Gly Lys Val Val Glu Ala Ser Ala 850 855 860
- Phe Gly Ile His Lys Gly Ser Ser Cys Asp Thr Val Ala Val Lys Met 865 870 875 880
- Leu Lys Glu Gly Ala Thr Ala Ser Glu Gln Arg Ala Leu Met Ser Glu 885 890 895
- Leu Lys Ile Leu Ile His Ile Gly Asn His Leu Asn Val Val Asn Leu 900 905 910
- Leu Gly Ala Cys Thr Lys Pro Gln Gly Pro Leu Met Val Ile Val Glu 915 920 925
- Phe Cys Lys Tyr Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys Arg Asp 930 935 940
- Ala Phe Ser Pro Cys Ala Glu Lys Ser Pro Glu Gln Arg Gly Arg Phe 945 950 955 960
- Arg Ala Met Val Glu Leu Ala Arg Leu Asp Arg Arg Arg Pro Gly Ser 965 970 975
- Ser Asp Arg Val Leu Phe Ala Arg Phe Ser Lys Thr Glu Gly Gly Ala 980 985 990
- Arg Arg Ala Ser Pro Asp Gln Glu Ala Glu Asp Leu Trp Leu Ser Pro
 995 1000 1005
- Leu Thr Met Glu Asp Leu Val Cys Tyr Ser Phe Gln Val Ala Arg Gly
 1010 1015 1020

Met Glu Phe Leu Ala Ser Arg Lys Cys Ile His Arg Asp Leu Ala Ala 025 1030 1035 1040

Arg Asn Ile Leu Leu Ser Glu Ser Asp Val Val Lys Ile Cys Asp Phe 1045 1050 1055

Gly Leu Ala Arg Asp Ile Tyr Lys Asp Pro Asp Tyr Val Arg Lys Gly
1060 1065 1070

Ser Ala Arg Leu Pro Leu Lys Trp Met Ala Pro Glu Ser Ile Phe Asp 1075 1080 1085

Lys Val Tyr Thr Thr Gln Ser Asp Val Trp Ser Phe Gly Val Leu Leu 1090 1095 1100

Trp Glu Ile Phe Ser Leu Gly Ala Ser Pro Tyr Pro Gly Val Gln Ile 105 1110 1115 1120

Asn Glu Glu Phe Cys Gln Arg Val Arg Asp Gly Thr Arg Met Arg Ala 1125 1130 1135

Pro Glu Leu Ala Thr Pro Ala Ile Arg His Ile Met Leu Asn Cys Trp 1140 1145 1150

Ser Gly Asp Pro Lys Ala Arg Pro Ala Phe Ser Asp Leu Val Glu Ile 1155 1160 1165

Leu Gly Asp Leu Leu Gln Gly Arg Gly Leu Gln Glu Glu Glu Val 1170 1175 1180

Cys Met Ala Pro Arg Ser Ser Gln Ser Ser Glu Glu Gly Ser Phe Ser 185 1190 1195 1200

Gln Val Ser Thr Met Ala Leu His Ile Ala Gln Ala Asp Ala Glu Asp 1205 1210 1215

Ser Pro Pro Ser Leu Gln Arg His Ser Leu Ala Ala Arg Tyr Tyr Asn 1220 1225 1230

Trp Val Ser Phe Pro Gly Cys Leu Ala Arg Gly Ala Glu Thr Arg Gly 1235 1240 1245

Ser Ser Arg Met Lys Thr Phe Glu Glu Phe Pro Met Thr Pro Thr Thr 1250 1255 1260

Tyr Lys Gly Ser Val Asp Asn Gln Thr Asp Ser Gly Met Val Leu Ala 265 1270 1275 1280

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Phe Arg

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